Access DB# 20937

SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name:		Examiner # :	Date:
Art Unit: Phone N	Number 30	Serial Numbe	r:
Mail Box and Bldg/Room Location	n:	_ Results Format Preferred	l (circle): PAPER DISK E-MAI
If more than one search is subm	nitted, please p	rioritize searches in orde	er of need.
Please provide a detailed statement of the Include the elected species or structures, I utility of the invention. Define any terms known. Please attach a copy of the cover	keywords, synonyn that may have a sp	ns, acronyms, and registry numb secial meaning. Give examples	ers, and combine with the concept or
Title of Invention:			
Inventors (please provide full names):			
Earliest Priority Filing Date:	· · ·		
For Sequence Searches Only Please inclu appropriate serial number.	ide all pertinent info	rmation (parent, child, divisional,	or issued patent numbers) along with the
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STAFF USE ONLY	Type of Searc		and cost where applicable
Searcher: M. Sm. Th	_ NA Sequence (#)		
Searcher Phone #:	_ AA Sequence (#)		
Searcher Location:	Structure (#)		
Date Searcher Picked Up:	_ Bibliographic _		
Date Completed: 8860	_ Litigation _	Lexis/Nexis	
Searcher Prep & Review Time: 15	_ Fulltext _	Sequence Systems	
Clerical Prep Time:	Patent Family	WWW/Internet	
	•	Other (cnecify)	

PTO-1590 (1-2000)

STIC-Biotech/ChemLib

From:

Elliott, George

Sent:

Tuesday, July 18, 2000 8:54

To:

STIC-Biotech/ChemLib

Cc: Subject: Nguyen, Qyang (AU1632)

FW: RUSH sequence search equest for 09/214124

Please rush.

Thanks,

George

Point of Contact: Mona Smith **Technical Info. Specialist** CM1 12C14 Tel: 308-3278

----Original Message-----

From: Sent:

Nguyen, Quang (AU1632) Tuesday, July 18, 2000 8:21 AM

To:

Elliott, George

Subject:

RUSH sequence search request for 09/214124

I would like to request a RUSH search request for the above application. Thanks.

For STIC-BIOTECH:

09/214124

Please search for nucleotide sequence that is homologous or identical to all or part of SEQ ID NO:1.

Please search for nucleotide sequence that is homologous or identical to SEQ ID NO:2 and to the following fragments of SEQ ID NO:2:

(a) Starting at nucleotide 1 and ending at nucleotide 578.

(b) Starting at nucleotide 265 and ending at nucleotide 578.

(c) Starting at nucleotide 452 and ending at nucleotide 578.

Against commercial, issued US application, interference databases. If there are questions regarding to this request, please contact me at 308-8339.

THANK YOU.